

FIG.I

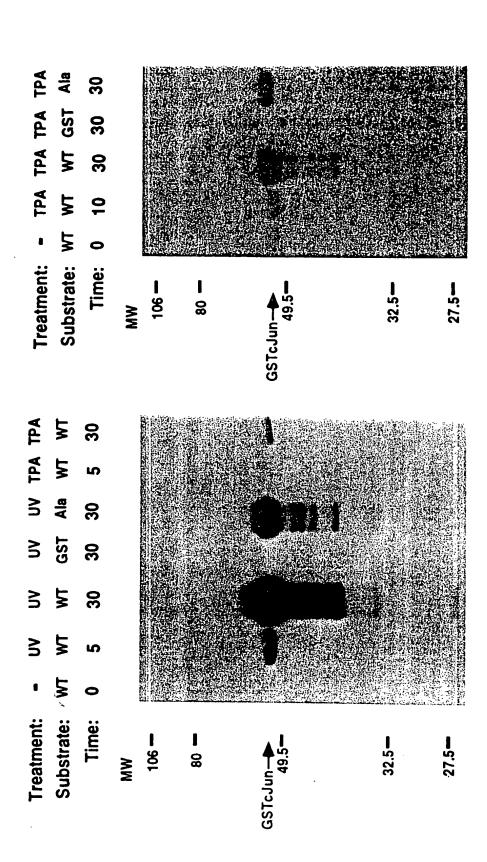
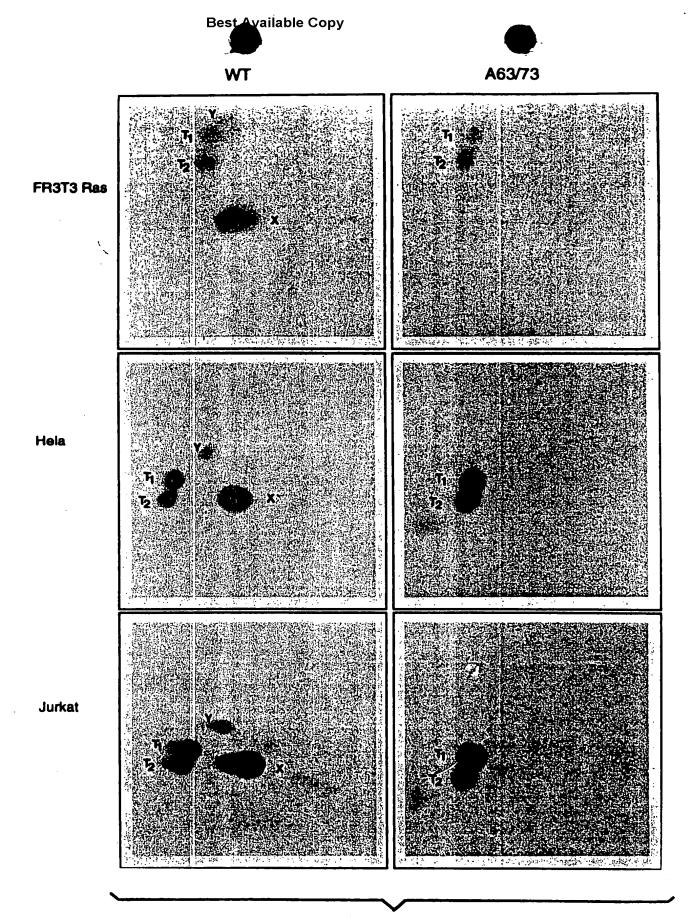


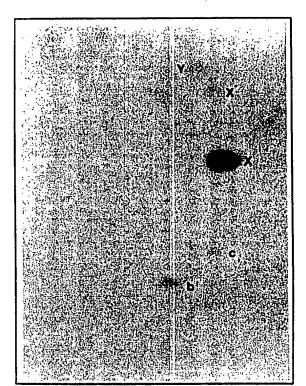
FIG.2B

F16.2A



FIĞ. 3A

In Vitro



In Vivo

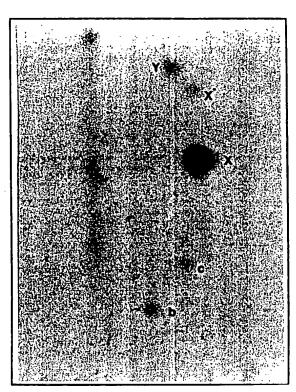
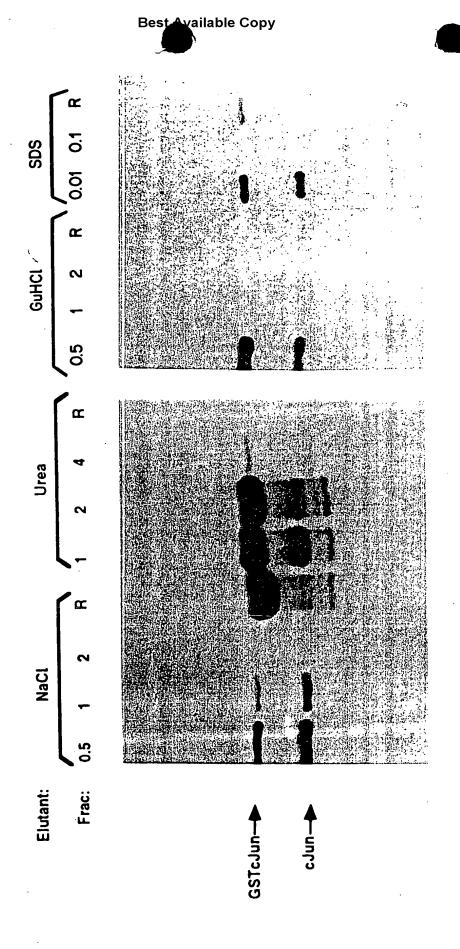


FIG. 3B



F16.4A

1 2 3 4 5

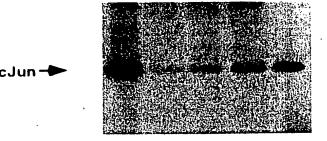
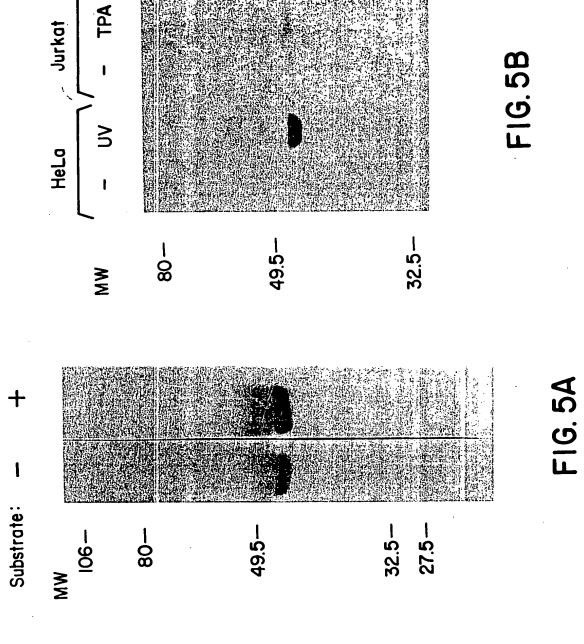


FIG. 4B



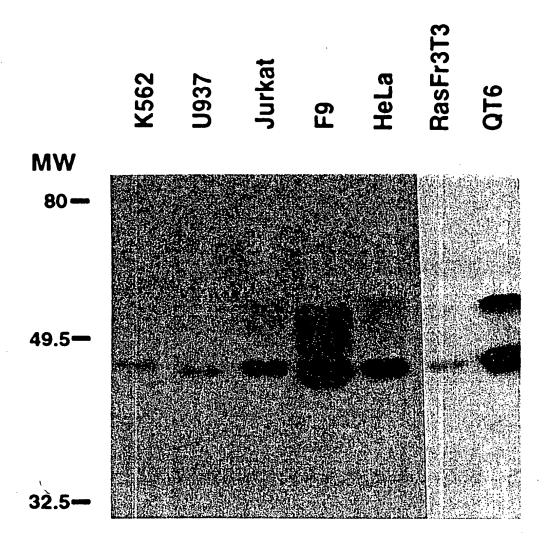


FIG.5C

	•							
GST	GSTcJ(1-223)	GSTcJ(11-223)	GSTcJ(22-223)	GSTcJ(33-223)	GSTcJ(43-223)	GSTcJ(56-223)	GSTcJ(1-93)	GSTcJ(1-79)
					のでは、中心では、神経のは、神経のなど、神経のなど、神経のなど、神経のなど、神経のなど、神経のなど、神経のなど、神経のなど、神経のなど、神経のなど、神経のなど、神経のなど、神経のなど、神経のなど、神経のなど			
CST	GSTcJ(1-223)	GSTcJ(11-223) GSTcJ(22-223)	GSTcJ(33-223)	GSTcJ(43-223)				

Protein Gel

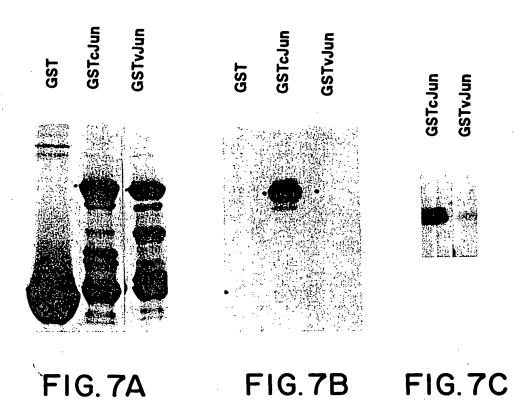
FIG.6A

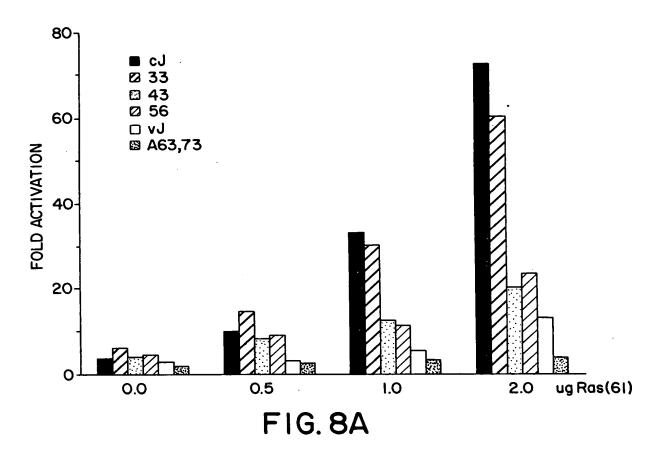
32<sub>P-Immobilized</sub> Substrate

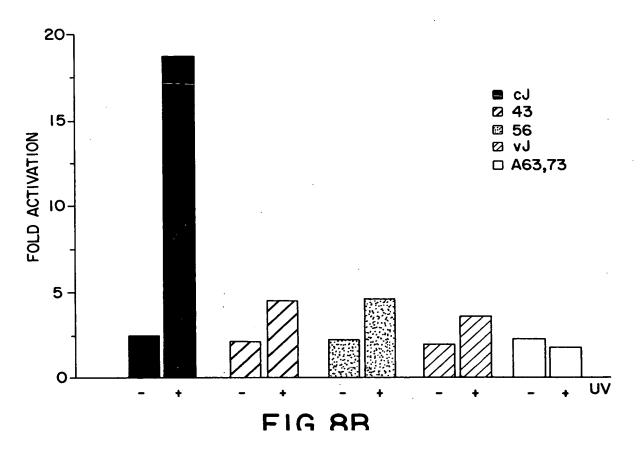
FIG.6B

32<sub>P-Exogenous</sub> Substrate

FIG.6C







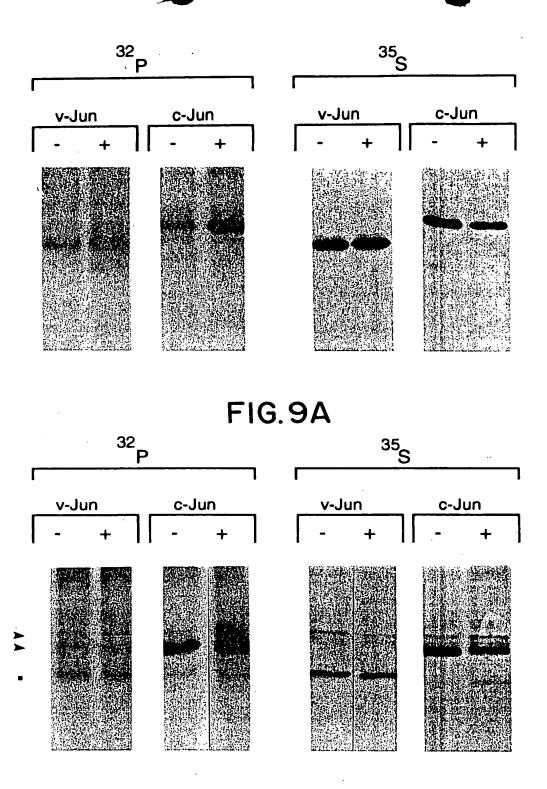


FIG.9B

	FIG.10A
561	AAA CAG AGC ATG ACC CTG ACC CTG GCC GAC CCA GTG GGG AGC CTG AAG Lys Gln Ser Met Thr Leu Asn Leu Ala Asp Pro Val Gly Ser Leu Lys 35
513	CTC CCC TCC GAG AGG GGA CCT TAT GGC TAC AGT AAC CCC AAG ATC CTG Leu Pro Ser Glu Arg Gly Pro Tyr Gly Tyr Ser Asn Pro Lys Ile Leu 20
765	GCA AAG ATG GAA ACG ACC TTC TAT GAC GAT GCC CTC AAC GCC TCG TTC Ala Lys Met Glu Thr Thr Phe Tyr Asp Asp Ala Leu Asn Ala Ser Phe 5
417	AACTIGIGCG CGCACGCCAA ACTAACCICA CGIGAAGIGA CGGACTGIIC I AIG ACT Met Ihr 1
360	STCAAAGGCT CCGGGGGGG CGGGTGTCCC CCGCTTGCCA CAGCCCTGTT GCGGCCCCGA
300	SAGGAGGGCG CACGGGGACG ACAGCCAGCG GGTGCGTGCG CTCTTAGAGA AACTTTCCCT
240	rcgacaagta agagtgcggg aggcatctta attaaccctg cgctccctgg agcagctggt
180	SCGCACGAAG AGCCGTCAGT GAGTGACCGC GACTTTTCAA AGCCGGGTAG GGCGGGGAG
120	STCCGCCGCA GAGCCGCTGC TCTGGGAAGT CAGTTCGCCT GCGGACTCCG AGGAACCGCT
09	BAATTCCGGG GCGCCAAGA CCCGCCGCCG GCCGGCCACT GCAGGGTCCG CACTGATCCG

CGG CAC CTC GGC GCC AAG AAC TCG GAC CTC ACC TGG GCC GAC GTC  Pro His Leu Arg Ala Lys Asn Ser Asp Leu Leu Thr Ser Pro Asp Val  GGG CTG CTC AAG CTG GCG TCG CGG GAG CTG GAG GCC CTG ATA ATC CAG  GLY Leu Leu Lys Leu Ala Ser Pro Glu Leu Glu Arg Leu Ile Ile Gln  Ser Ser Asn Gly His Ile Thr Thr Thr Thr Pro Thr Gln Pre Leu  Ss Ser Ser Asn GGC CAC ACC ACC ACC CGC ACC CAG TCG  GGG CGC CAC ATC ACC ACC ACC CGC ACC CAG TCG  TCC CAG AAC GGG CAC ATC ACC ACC ACC CCC ACC CAG TCG  Ser Ser Asn Gly His Ile Thr Thr Thr Thr Tro Thr Gln Pre Leu  GGG CGC CAC ACC ACC ACC CAC ACC CGC ACC CAG TCG  TGC CCC AAG AAC GTG ACA CAC ACC CAG ACC CCC ACC CAG TCC  Cys Pro Lys Asn Val Thr Asp Glu Glu Gly Pre Ala Glu Gly Pre  TGC CCC AAG ACC GCC CAC ACC CAC ACC CAC ACC CCC C	609	657	705	753	801	849	897
CGC CGC CAGC AAC ACC CCC CGC CGC CGC CGC							
CAG CTC CGC GCC AAG AAC TCG GAC CTC CTC ACC TCG CCC His Leu Lu Thr Ser Pro 60 CTC AAG ATA Leu Leu Lys Leu Ala Ser Pro Glu Leu Glu Arg Leu Ile 80 AGC AAC GGC CTC ACC CAG ATA AGC AAC ACC ACC ACC CCC ACC CAG ACC ACC	GTG Val	CAG Gln	CIG	TTC Phe	GTC Val 130	GCG Ala	
CCC CCC CCC CAC AAC TCG GAC CTC CTC ACC TCC His Leu Arg Ala Lys Asn Ser Asp Leu Leu Thr Ser CTG CTC AAC CTG GCG TCG CCC GAG CTG GAG CGC CTG Leu Leu Lys Leu Ala Ser Pro Glu Leu Glu Arg Leu ACC AAC GCG CAC ATC ACC ACC ACC CCC ACC Ser Asn Gly His Ile Thr Thr Thr Pro Thr Pro Thr 85 CCC AAG AAC GTG ACA GAT GAG CAG GGG TTC GCC Pro Lys Asn Val Thr Asp Glu Gln Glu Gly Phe Ala 100 CGC GCC CTG GCC GAA CTG CAC AGC CAG AAC ACG ATG ALa Leu Ala Glu Leu His Ser Gln Asn Thr Leu ATG GCG CCG CAG CCG GTC AAC GGG GCG ATG CGC GCG CTG GCG GGG GGG GGG GTG Ser Ala Ala Gln Pro Val Asn Gly Ala Gly Met Val 135 GCC TCG GTG GCA GGG GGC AGC GGC GTC Ala Ser Val Ala Gly Ser Gly Ser Gly Gly Phe 150 Ala Ser Val Ala Gly Ser Gly Ser Gly Gly Phe	GAC Asp 65	ATC Ile	TTC	660 61y	AGC	CCC Pro 145	GCC Ala
CAC CTC CGC GCC AAG AAC TCG GAC CTC CTC ACC His Leu Arg Ala Lys Asn Ser Asp Leu Leu Thr CTG CTG GCG GCG GCG GCG GCG GCG GCG GCG	CCC	ATA Ile 80	CAG Gln	GAG Glu	CCC	GCT	AGC Ser 160
CTC CGC GCC AAG AAG TCG GAC CTC CTC His Leu Arg Ala Lys Asn Ser Asp Leu Leu Luys Leu Ala Ser Pro Glu Leu Glu Leu Lys Leu Ala Ser Pro Glu Leu Glu Leu Lys Leu Ala Ser Pro Glu Leu Glu Leu Lys Leu Ala Ser Pro Glu Leu Glu Leu Lys Asn GTG ACG ACG ACG ACG ACG ACG AGG AGG AGG GCG AGG AG	TCG	CTG Leu	ACC Thr 95	GCC Ala	CTG	GTG Val	
CAC CTC GGC GCC AAG ASG TCG GAC CTC His Leu Arg Ala Lys Asn Ser Asp Leu 555  CTG CTC AAG CTG GCG TCG CCC GAG CTG Leu Lys Leu Ala Ser Pro Glu Leu Lys Leu Ala Ser Pro Glu Leu Lys Leu Ala Ser Pro Glu Leu Lys Leu Ala GTG ACC ACC ACG CCG Ser Asn Gly His Ile Thr Thr Thr Thr Pro Pro Lys Asn Val Thr Asp Glu Glu Glu 100  CGC GCC CTG GCC GAA CTG CAC AGG CAG ACG GCC ATG ALa Glu Leu His Ser Gln 120  TCG GCC CTG GCG CAG GCG GCG GCA Ser Ala Ala Gln Pro Val Asn Gly Ala Ser Ala Ala Gln Pro Val Asn Gly Ala Ser Val Ala Gly Gly Gly Ser Iso	ACC Thr	CGC Arg	CCC	TTC Phe 110	ACG Thr	ATG	660 61y
CAC CTC CGC GCC AAG AAC TCG GAC His Leu Arg Ala Lys Asn Ser Asp 55 CTG CTC AAG CTG GCG TCG CCC GAG Leu Leu Lys Leu Ala Ser Pro Glu 70 Ser Asn Gly His Ile Thr Thr Thr Thr Thr 100 CCC AAG AAC GTG ACA GAT GAG CAG Pro Lys Asn Val Thr Asp Glu Gln 100 TCG GCC CTG GCC GAA CTG CAC AGG Arg Ala Leu Ala Glu Leu His Ser GCG Ser Ala Ala Gln Pro Val Asn Gly Ser Ala Ser Val Ala Gly Ser Gly Ser Val Ala Ser Val Ala Gly Gly Ser Gly Ala Ser Val Ala Gly Gly Ser Gly Ala Ser Val Ala Gly Gly Ser Gly Iso	CIC	GAG Glu	ACC	666 61y	AAC Asn 125	GGC Gly	660 61y
CAC CTC CGC GCC AAG AAC THis Leu Arg Ala Lys Asn S 55 CTG CTC GCG TCG CTG GCG TCG CTG AAC GTG GTG GTG GTG GTG GTG GTG GTG GTG GT	CTC Leu 60	CTG	CCG	GAG	CAG Gln		AGC Ser
CAC CTC CGC GCC AAG AAC THis Leu Arg Ala Lys Asn S 55 CTG CTC GCG TCG CTG GCG TCG CTG AAC GTG GTG GTG GTG GTG GTG GTG GTG GTG GT	GAC Asp	GAG Glu 75	ACG Thr	CAG Gln	AGC Ser	GGG	660 61y 155 <b>G. IC</b>
CAC CTC CGC GCC AAG His Leu Arg Ala Lys CTG CTC AAG CTG GCG Leu Leu Lys Leu Ala AGC AAC GGG CAC ATC Ser Asn Gly His Ile 85 CCC AAG AAC GTG ACA Pro Lys Asn Val Thr 100 CGC GCC CTG GCC GAA Arg Ala Leu Ala Glu 120 TCG GCG CGG CAG CCG Ser Ala Ala Gln Pro GCC TCG GTG GCA GCC TCG GTG GCA Ala Ser Val Ala Gly Ala Ser Val Ala Gly Ala Ser Val Ala Gly	TCG	CCC Pro	ACC Thr 90	GAG Glu	CAC	AAC Asn	4, 01
CAC CTC CGC GCC A His Leu Arg Ala I Leu Leu Lys Leu A Ser Asn Gly His I NO CCC AAG AAC GTG A Ser Asn Gly His I 100 CGC GCC CTG GCC G Arg Ala Leu Ala Gln F Ser Ala Ala Gln F 135 GCC TCG GTG GCA G Ser Ala Ala Gln F 135 GCC TCG GTG GCA G Ala Ser Val Ala G Ala Ser Val Ala G	AAC Asn	TCG Ser	ACC	GAT Asp 105	CTG	GTC Val	
CAC CTC CGC GCC His Leu Arg Ala 55 CTG CTC AAG CTG Leu Leu Lys Leu 85 CCC AAG AAC GTG Pro Lys Asn Val 100 CGC GCC CTG GCC Arg Ala Leu Ala 135 GCC TCG GCG CAG Ser Ala Ala Gln 135 GCC TCG GTG GCA Ala Ser Val Ala 150	AAG Lys	GCG Ala	ATC	ACA	GAA Glu 120	CCG	666 61y
CAC CTC Lis Leu CTG CTC Leu Leu Ser Asn 85 CCC AAG Pro Lys 100 CGC GCC Arg Ala TCG GCG Ser Ala GCC TCG Ala Ala Ser	$\mathcal{O} \vdash \mathcal{O}$	CTG Leu	C) W	⊟ ત	$\mathcal{O}$	CAG Gln 135	GCA Ala
CAC CTG Leu CGC CGC AFBO Ser CGC AFB	CGC	AAG Lys 70	666 G1y	AAC Asn	CTG Leu	GCG Ala	Han
	CIC Leu	CTC	AAC Asn 85	AAG Lys	GCC	$\circ$	ပေၿ
CCG Pro Pro GGG G1y Cys Cys Cys GTG Thr Val	CAC	CTG	மும	CCC Pro 100	CGC Arg	ပေၿ	0 H
	CCG	O H	ပေၿ	TGC Cys	GTG Val 115	ACG Thr	GTA Val

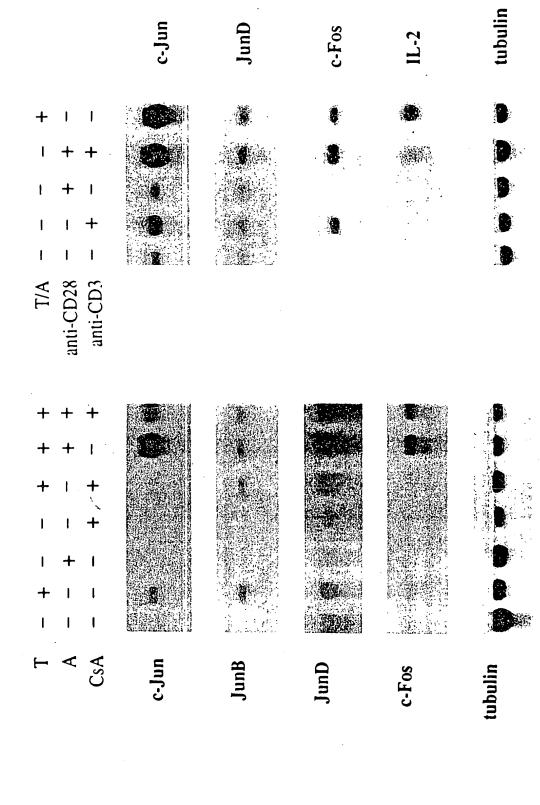
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AAC CCA Asn Pro	GCC GGC Ala Gly	CAC CAC His His 210	CTG AAG Leu Lys 225	CCC CTG Pro Leu	AGG AAG Arg Lys	AAG CTG Lys Leu
AAC TTC Asn Phe 175	GGC GCG Gly Ala	CCG CCG Pro Pro	CAG GCC Gln Ala	ACA CCG Thr Pro	GCG GAG Ala Glu A	AAA AGG L Lys Arg
CTC AGC Leu Ser	TCC TAC Ser Tyr 190	CAG CAG Gln Gln 205	CGG CTG Arg Leu	GGC GAG Gly Gly G	ATC AAG	TGC CGA Cys Arg 270
GCA AAC Ala Asn	cg ccc la Pro	CAG CAG Gln Gln	CAC CCG ( His Pro / 220	ATG CCC ( Met Pro ( 235	GAG CGC / Glu Arg ]	TCG AAG 1 Ser Lys (
C TAC 1 TYF 170	GC GGG G 1y Gly A 85	c CAG o Gln	G CAG 1 Gln	c GAG o Glu	c CAG r Gln 250	r GCC a Ala 5
G CCG GTO Pro Va	C GGC GC r Gly G]	GAA CC	G CCC GT t Pro Va 5	A GTG CC e Val Pr	G GAG TCO t Glu Ser	ATC GC File Al
GAG CCG Glu Pro	AGC AG( Ser Se)	CCC GCG Pro Ala	CAG AT Gln Me	CAG AT Gln Il 230	GAC ATC Asp Met	AAC CGC Asn Arg
CAC AGC His Ser 165	GCG CTG Ala Leu 180	GCC TTT Ala Phe	CCC CAG Pro Gln	GAG CCT Glu Pro	CCC ATC Pro Ile 245	ATG AGG Met Arg 260
CTG	66C G1y	CTG Leu 195	CTG	GAG	TCC	CGC

F1G.10C

	-		(			•			
1281	1329	1377	1424	1484	1544	1604	1664	1724	
GAG AGA ATC GCC CGG CTG GAG GAA AAA GTG AAA ACC TTG AAA GCT CAG Glu Arg Ile Ala Arg Leu Glu Glu Lys Val Lys Thr Leu Lys Ala Gln 275	AAC TCG GAG CTG GCG TCG ACG GCC AAC ATG CTC AGG GAA CAG GTC GCA Asn Ser Glu Leu Ala Ser Thr Ala Asn Met Leu Arg Glu Gln Val Ala 300	CAG CTT AAA CAC AAA GTC ATG AAC CAC GTT AAC AGT GGG TGC CAA CTC Gln Leu Lys His Lys Val Met Asn His Val Asn Ser Gly Cys Gln Leu 310	ATC CTA ACG CAG TTG CAA ACA TTT TGAAGAGA CCGTCGGGGG Ile Leu Thr Gln Gln Leu Gln Thr Phe 325	CIGAGGGGCA ACGAAGAAA AAAATAACAC AGAGAGACAG ACTIGAGAAC ITGACAAGIT	GCGACGGAGA GAAAAAAAA GTGTCCGAGA ACTAAAGCCA AGGGTATCCA AGTTGGACTG	GGTICGGICT GACGCCCCC CCAGIGIGCA CGAGIGGGAA CCACCIGGIC GCGCCTCCC	IIGGCGICGA GCCAGGGAGC GGCCGCCIGG GGGCIGCCCC GCTIIGCGGA CGGGCIGICC	CCGCGCGAAC GGAACGTTGG ACTTTCGTTA ACATTGACCA AGAACTGCAT GGACCTAACA	FIG.10D

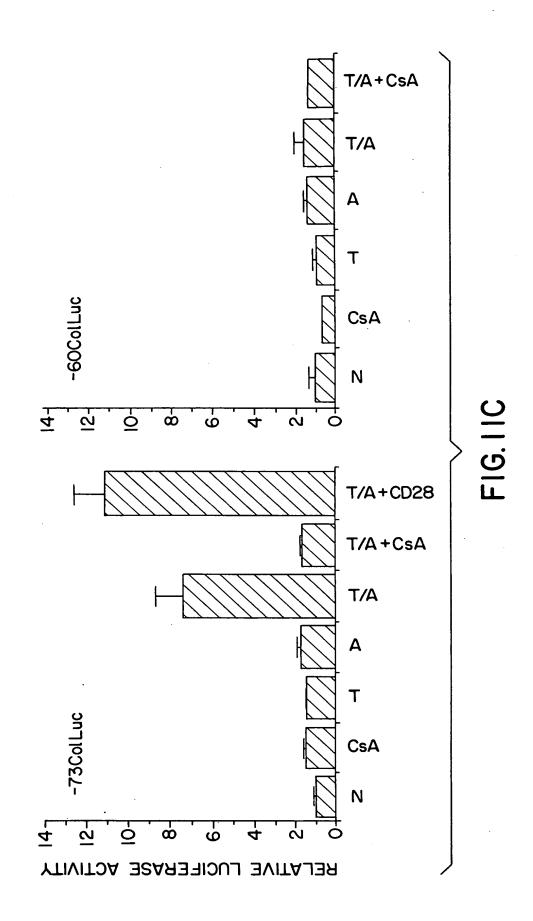
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TGTAGATTGC	IGTAGATIGC TICTGTAGTA CTCCTTAAGA ACACAAAGCG GGGGGAGGGT TGGGGAGGGG	CTCCTTAAGA	ACACAAAGCG	GGGGGAGGGT	TGGGGAGGGG	1847
CGGCAGGAGG	CGGCAGGAGG GAGGTITGIG AGAGGGAGGC IGAGCCIACA GAIGAACICI IICIGGCCIG	AGAGCGAGGC	TGAGCCTACA	GATGAACTCT	TICIGGCCIG	1907
CTTTCGTTAA	CTTTCGTTAA CTGTGTATGT ACATATATA ATTTTTAAT TTGATTAAAG CTGATTACTG	ACATATATAT	ATTTTTAAT	TTGATTAAAG	CIGAITACIG	196
TCAATAAACA	TCAATAAACA GCTTCATGCC TTTGTAAGTT ATTTCTTGTT TGTTTG GGATCCTGCC	TTTGTAAGTT	ATTTCTTGTT	TGTTTGTTTG	GGATCCTGCC	205
CAGTGTTGTT	CAGTGTTGTT TGTAAATAAG AGATTTGGAG CACTCTGAGT TTACCATTTG TAATAAAGTA	AGATTTGGAG	CACTCTGAGT	TTACCATTTG	TAATAAAGTA	208
ىدەدە دىنىدىدىدىد لىد	F-F					209

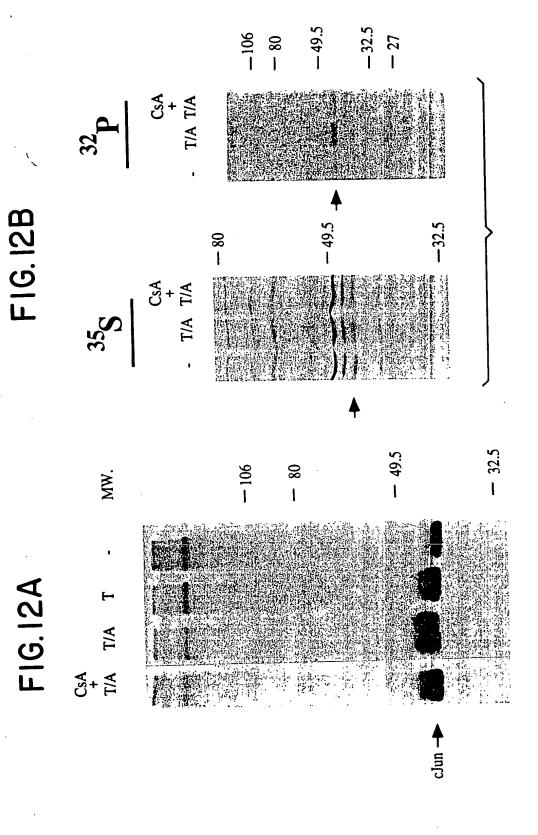
F16.10E

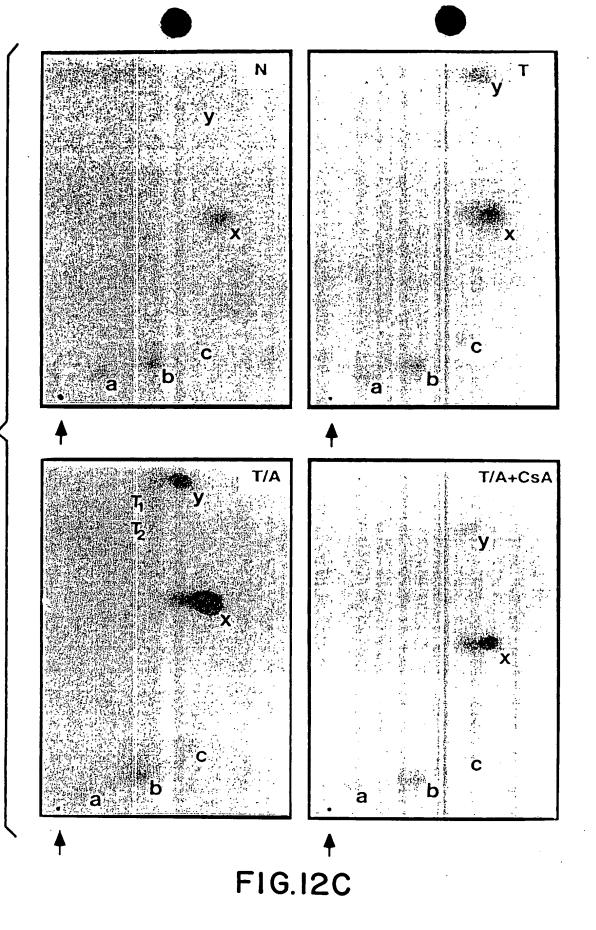


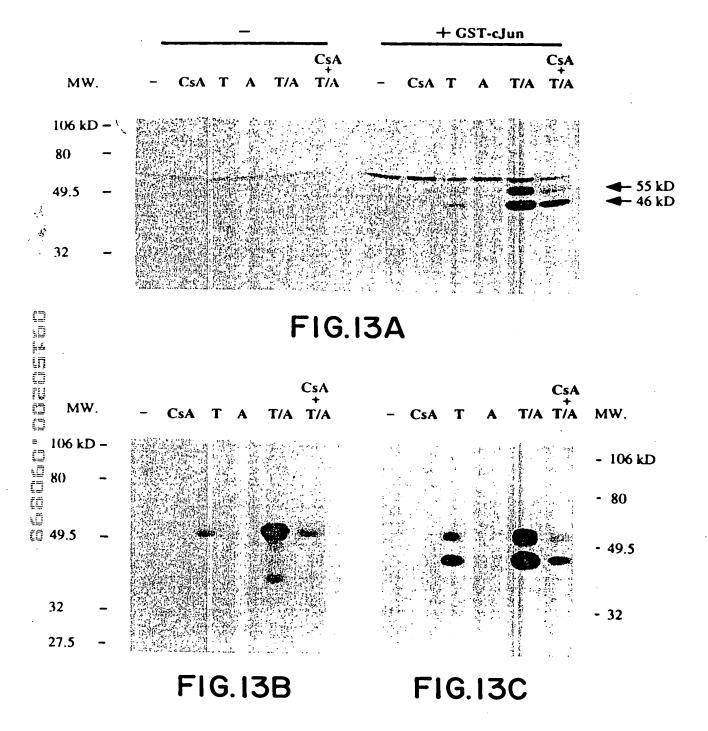
F16.11B

FIG.11A









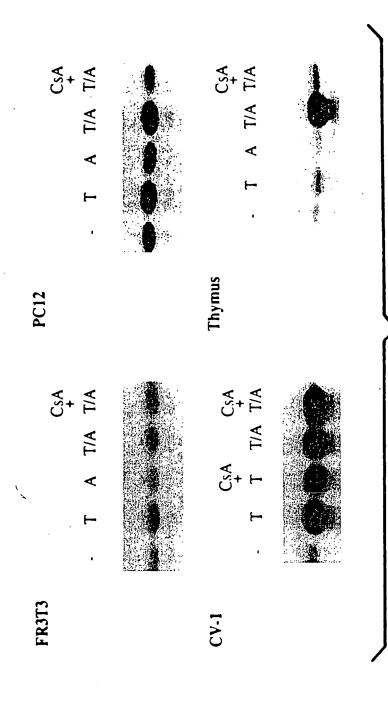


FIG.14

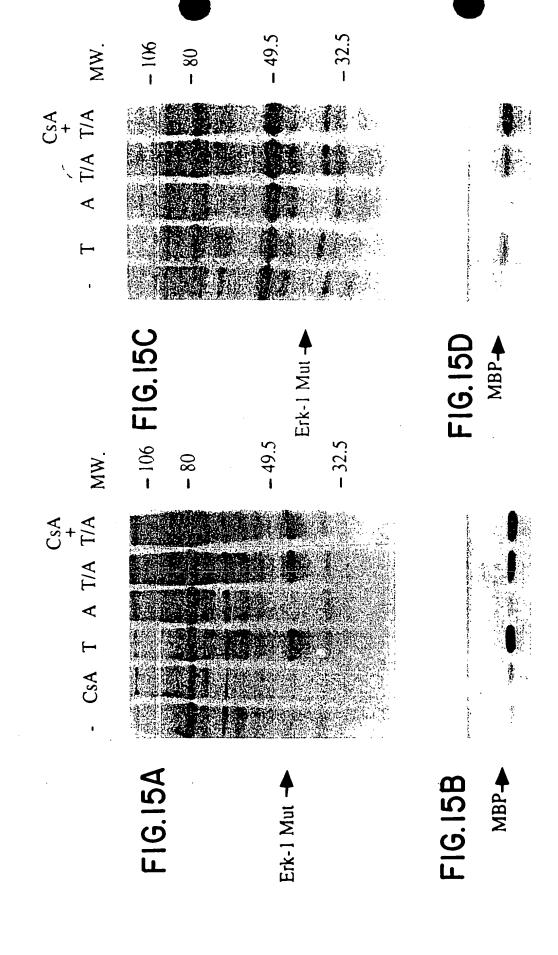


FIG.16A

FIG. 16B

